

Acta Crystallographica Section E

Structure Reports

Online

ISSN 1600-5368

Ethyl 2-[[4-(pyridin-4-yl)pyrimidin-2-yl]-sulfanyl]acetate

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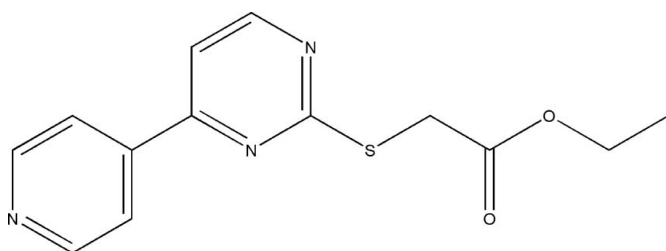
Received 31 January 2011; accepted 4 February 2011

Key indicators: single-crystal X-ray study; $T = 291$ K; mean $\sigma(\text{C}-\text{C}) = 0.003$ Å; R factor = 0.038; wR factor = 0.115; data-to-parameter ratio = 17.5.

In the title molecule, $\text{C}_{13}\text{H}_{13}\text{N}_3\text{O}_2\text{S}$, the pyridine and pyrimidine rings form a dihedral angle of 3.8 (1°). The crystal packing exhibits weak intermolecular $\text{C}-\text{H}\cdots\text{O}$ hydrogen bonds.

Related literature

For details of the synthesis and general background to the rational design and assembly of coordination polymers with thioethers, see: Dong *et al.* (2008, 2009). For the crystal structures of coordination complexes with related ligands, see: Du *et al.* (2004); Zhu *et al.* (2009).



Experimental

Crystal data

 $\text{C}_{13}\text{H}_{13}\text{N}_3\text{O}_2\text{S}$
 $M_r = 275.33$

 Triclinic, $P\bar{1}$
 $a = 8.6579$ (8) Å

 $b = 9.7394$ (9) Å
 $c = 9.9188$ (8) Å
 $\alpha = 62.661$ (6°)
 $\beta = 71.416$ (5°)
 $\gamma = 65.024$ (6°)
 $V = 665.35$ (10) Å³
 $Z = 2$
 Mo $K\alpha$ radiation
 $\mu = 0.25$ mm⁻¹
 $T = 291$ K
 $0.32 \times 0.24 \times 0.18$ mm

Data collection

 Bruker SMART CCD area-detector diffractometer
 Absorption correction: multi-scan (SADABS; Bruker, 2000)
 $T_{\min} = 0.917$, $T_{\max} = 0.966$

 11760 measured reflections
 3021 independent reflections
 2387 reflections with $I > 2\sigma(I)$
 $R_{\text{int}} = 0.032$

Refinement

 $R[F^2 > 2\sigma(F^2)] = 0.038$
 $wR(F^2) = 0.115$
 $S = 1.04$
 3021 reflections

 173 parameters
 H-atom parameters constrained
 $\Delta\rho_{\text{max}} = 0.24$ e Å⁻³
 $\Delta\rho_{\text{min}} = -0.20$ e Å⁻³

Table 1

Hydrogen-bond geometry (Å, °).

$D-\text{H}\cdots A$	$D-\text{H}$	$\text{H}\cdots A$	$D\cdots A$	$D-\text{H}\cdots A$
$\text{C3}-\text{H3}\cdots\text{O1}^{\text{i}}$	0.93	2.52	3.383 (2)	154
$\text{C7}-\text{H7}\cdots\text{O1}^{\text{ii}}$	0.93	2.61	3.373 (2)	140

Symmetry codes: (i) $-x + 1, -y, -z + 2$; (ii) $x, y, z + 1$.

Data collection: SMART (Bruker, 2000); cell refinement: SAINT (Bruker, 2000); data reduction: SAINT; program(s) used to solve structure: SHELXTL (Sheldrick, 2008); program(s) used to refine structure: SHELXTL; molecular graphics: SHELXTL; software used to prepare material for publication: SHELXTL.

The author is indebted to the National Natural Science Foundation of China (grant No. 20871039) for financial support.

Supplementary data and figures for this paper are available from the IUCr electronic archives (Reference: CV5047).

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supporting information

Acta Cryst. (2011). E67, o690 [doi:10.1107/S1600536811004272]

Ethyl 2-[[4-(pyridin-4-yl)pyrimidin-2-yl]sulfanyl]acetate**Chuan-Hu Wang****S1. Comment**

Remarkable attention has been paid to the rational design and assembly of new coordination polymers with thioethers (Dong *et al.*, 2008; 2009; Du *et al.*, 2004; Zhu *et al.*, 2009). Herewith we report the synthesis and crystal structure of the title compound (I) - a new derivative of 4-(4-pyridinyl)pyrimidine-2-thiol.

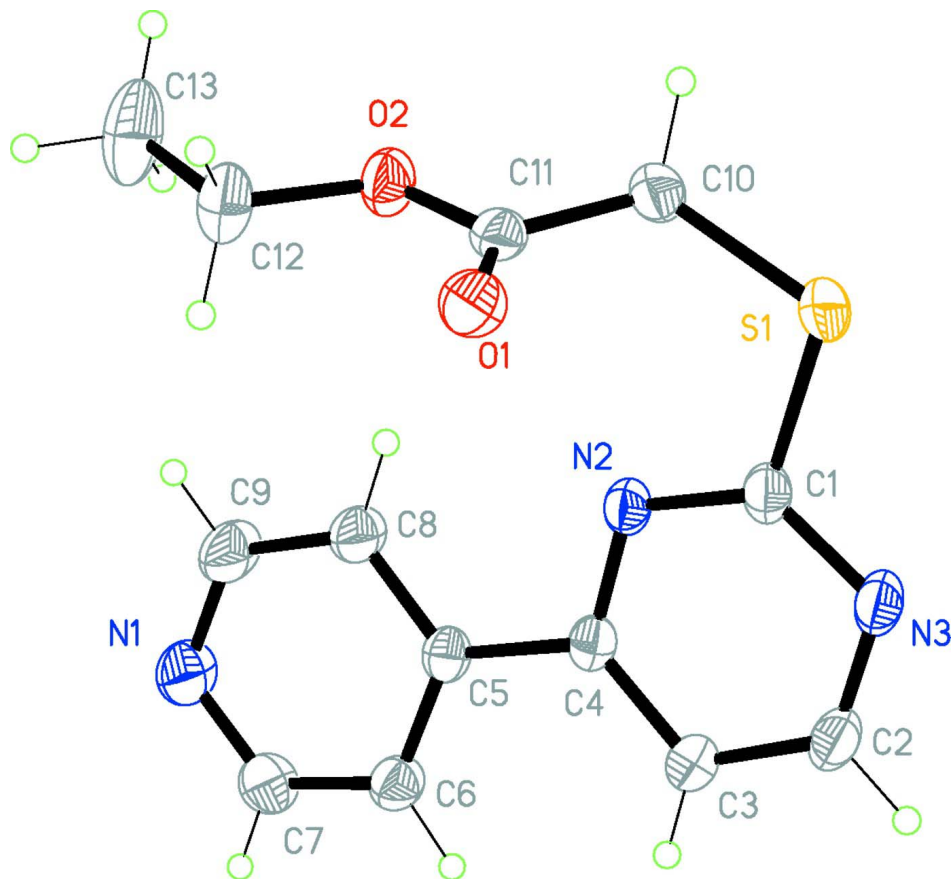
In (I) (Fig. 1), the pyridine and pyrimidine rings form a dihedral angle of 3.8 (1)°. The crystal packing exhibits weak intermolecular C—H···O hydrogen bonds (Table 1).

S2. Experimental

All solvents and chemicals were of analytical grade and were used without further purification. The title compound was prepared by similar procedure reported in the literature (Dong *et al.*, 2008; 2009), To a solution of 4-(4-pyridinyl)pyrimidine-2-thiol (3.78 g, 20 mmol) and sodium hydroxide (0.80 g, 20 mmol) in dry ethanol (100 ml), ethyl 2-bromoacetate (3.34 g, 20 mmol) in CCl₄ (30 ml) was added. The mixture was stirred and refluxed for 8 h. After cooling, precipitates were filtered, washed by water and ethanol, and dried in vacuum. Single crystals suitable for X-ray diffraction were grown from methanol solution by slow evaporation in air at room temperature.

S3. Refinement

All H atoms were geometrically positioned (C—H 0.93–0.97 Å) and refined as riding, with $U_{\text{iso}}(\text{H})=1.2-1.5 U_{\text{eq}}$ of the parent atom.

**Figure 1**

The structure of the title compound, showing 30% probability displacement ellipsoids and the atomic numbering.

Ethyl 2-[[4-(pyridin-4-yl)pyrimidin-2-yl]sulfanyl]acetate

Crystal data

$C_{13}H_{13}N_3O_2S$

$M_r = 275.33$

Triclinic, $P\bar{1}$

Hall symbol: -P 1

$a = 8.6579$ (8) Å

$b = 9.7394$ (9) Å

$c = 9.9188$ (8) Å

$\alpha = 62.661$ (6)°

$\beta = 71.416$ (5)°

$\gamma = 65.024$ (6)°

$V = 665.35$ (10) Å³

$Z = 2$

$F(000) = 288.0$

$D_x = 1.374$ Mg m⁻³

Mo $K\alpha$ radiation, $\lambda = 0.71073$ Å

Cell parameters from 3021 reflections

$\theta = 2.3$ – 27.5 °

$\mu = 0.25$ mm⁻¹

$T = 291$ K

Block, colorless

$0.32 \times 0.24 \times 0.18$ mm

Data collection

Bruker SMART CCD area-detector
diffractometer

Radiation source: fine-focus sealed tube

Graphite monochromator

φ and ω scans

Absorption correction: multi-scan
(*SADABS*; Bruker, 2000)

$T_{\min} = 0.917$, $T_{\max} = 0.966$

11760 measured reflections

3021 independent reflections

2387 reflections with $I > 2\sigma(I)$

$R_{\text{int}} = 0.032$
 $\theta_{\text{max}} = 27.5^\circ$, $\theta_{\text{min}} = 2.3^\circ$
 $h = -11 \rightarrow 10$

$k = -12 \rightarrow 12$
 $l = -12 \rightarrow 12$

Refinement

Refinement on F^2
 Least-squares matrix: full
 $R[F^2 > 2\sigma(F^2)] = 0.038$
 $wR(F^2) = 0.115$
 $S = 1.04$
 3021 reflections
 173 parameters
 0 restraints
 Primary atom site location: structure-invariant
 direct methods

Secondary atom site location: difference Fourier
 map
 Hydrogen site location: inferred from
 neighbouring sites
 H-atom parameters constrained
 $w = 1/[\sigma^2(F_o^2) + (0.0658P)^2 + 0.067P]$
 where $P = (F_o^2 + 2F_c^2)/3$
 $(\Delta/\sigma)_{\text{max}} < 0.001$
 $\Delta\rho_{\text{max}} = 0.24 \text{ e } \text{\AA}^{-3}$
 $\Delta\rho_{\text{min}} = -0.20 \text{ e } \text{\AA}^{-3}$

Special details

Experimental. The structure was solved by direct methods (Bruker, 2000) and successive difference Fourier syntheses.

Geometry. All e.s.d.'s (except the e.s.d. in the dihedral angle between two l.s. planes) are estimated using the full covariance matrix. The cell e.s.d.'s are taken into account individually in the estimation of e.s.d.'s in distances, angles and torsion angles; correlations between e.s.d.'s in cell parameters are only used when they are defined by crystal symmetry. An approximate (isotropic) treatment of cell e.s.d.'s is used for estimating e.s.d.'s involving l.s. planes.

Refinement. Refinement of F^2 against ALL reflections. The weighted R -factor wR and goodness of fit S are based on F^2 , conventional R -factors R are based on F , with F set to zero for negative F^2 . The threshold expression of $F^2 > \sigma(F^2)$ is used only for calculating R -factors(gt) *etc.* and is not relevant to the choice of reflections for refinement. R -factors based on F^2 are statistically about twice as large as those based on F , and R -factors based on ALL data will be even larger.

Fractional atomic coordinates and isotropic or equivalent isotropic displacement parameters (\AA^2)

	<i>x</i>	<i>y</i>	<i>z</i>	$U_{\text{iso}}^*/U_{\text{eq}}$
C1	0.21141 (18)	0.00339 (18)	0.89933 (17)	0.0411 (3)
C2	0.3174 (2)	-0.21822 (19)	1.10105 (19)	0.0496 (4)
H2	0.3517	-0.3308	1.1555	0.060*
C3	0.3336 (2)	-0.12042 (18)	1.15846 (18)	0.0459 (4)
H3	0.3793	-0.1652	1.2484	0.055*
C4	0.27871 (17)	0.04721 (17)	1.07653 (16)	0.0376 (3)
C5	0.28647 (18)	0.16653 (17)	1.12645 (16)	0.0388 (3)
C6	0.3582 (2)	0.1176 (2)	1.25464 (18)	0.0478 (4)
H6	0.4018	0.0070	1.3141	0.057*
C7	0.3641 (2)	0.2347 (2)	1.2930 (2)	0.0568 (4)
H7	0.4126	0.1991	1.3794	0.068*
C8	0.2241 (2)	0.3332 (2)	1.04370 (19)	0.0525 (4)
H8	0.1740	0.3725	0.9573	0.063*
C9	0.2372 (3)	0.4407 (2)	1.0913 (2)	0.0660 (5)
H9	0.1960	0.5521	1.0335	0.079*
C10	0.0836 (2)	0.2913 (2)	0.6641 (2)	0.0515 (4)
H10A	0.0315	0.3439	0.5718	0.062*
H10B	-0.0042	0.3227	0.7439	0.062*
C11	0.2273 (2)	0.35667 (18)	0.63290 (16)	0.0422 (3)
C12	0.2788 (2)	0.5879 (2)	0.6083 (2)	0.0620 (5)
H12A	0.3426	0.6012	0.5056	0.074*

H12B	0.3603	0.5258	0.6805	0.074*
C13	0.1774 (3)	0.7498 (3)	0.6187 (3)	0.0927 (8)
H13A	0.1014	0.8124	0.5432	0.139*
H13B	0.2542	0.8066	0.6002	0.139*
H13C	0.1108	0.7354	0.7193	0.139*
N1	0.3051 (2)	0.39441 (19)	1.21466 (18)	0.0644 (4)
N2	0.21695 (15)	0.11055 (14)	0.94434 (14)	0.0392 (3)
N3	0.25523 (17)	-0.15909 (16)	0.97173 (16)	0.0482 (3)
O1	0.37818 (15)	0.29263 (14)	0.59842 (13)	0.0516 (3)
O2	0.15904 (14)	0.50272 (13)	0.64426 (13)	0.0512 (3)
S1	0.14603 (6)	0.07492 (5)	0.72151 (5)	0.05259 (16)

Atomic displacement parameters (Å²)

	U^{11}	U^{22}	U^{33}	U^{12}	U^{13}	U^{23}
C1	0.0408 (8)	0.0366 (8)	0.0520 (8)	-0.0153 (6)	-0.0068 (6)	-0.0197 (6)
C2	0.0542 (9)	0.0320 (8)	0.0601 (10)	-0.0176 (7)	-0.0095 (7)	-0.0119 (7)
C3	0.0514 (9)	0.0366 (8)	0.0489 (8)	-0.0168 (7)	-0.0111 (7)	-0.0113 (7)
C4	0.0365 (7)	0.0348 (7)	0.0428 (7)	-0.0142 (6)	-0.0023 (6)	-0.0159 (6)
C5	0.0402 (8)	0.0373 (8)	0.0415 (7)	-0.0142 (6)	-0.0015 (6)	-0.0188 (6)
C6	0.0573 (9)	0.0410 (8)	0.0463 (8)	-0.0143 (7)	-0.0124 (7)	-0.0162 (7)
C7	0.0713 (11)	0.0575 (11)	0.0533 (9)	-0.0186 (9)	-0.0167 (8)	-0.0282 (8)
C8	0.0712 (11)	0.0398 (9)	0.0521 (9)	-0.0135 (8)	-0.0201 (8)	-0.0192 (7)
C9	0.1000 (15)	0.0390 (9)	0.0678 (11)	-0.0152 (9)	-0.0285 (10)	-0.0238 (8)
C10	0.0549 (9)	0.0403 (8)	0.0677 (10)	-0.0094 (7)	-0.0271 (8)	-0.0213 (8)
C11	0.0533 (9)	0.0353 (8)	0.0405 (7)	-0.0108 (7)	-0.0175 (6)	-0.0130 (6)
C12	0.0637 (11)	0.0517 (10)	0.0791 (12)	-0.0294 (9)	-0.0024 (9)	-0.0277 (9)
C13	0.0989 (17)	0.0653 (14)	0.133 (2)	-0.0416 (13)	0.0127 (15)	-0.0589 (15)
N1	0.0889 (11)	0.0530 (9)	0.0664 (9)	-0.0200 (8)	-0.0205 (8)	-0.0321 (8)
N2	0.0434 (7)	0.0334 (6)	0.0460 (7)	-0.0143 (5)	-0.0076 (5)	-0.0172 (5)
N3	0.0534 (8)	0.0346 (7)	0.0629 (8)	-0.0176 (6)	-0.0102 (6)	-0.0199 (6)
O1	0.0513 (7)	0.0473 (7)	0.0572 (7)	-0.0104 (5)	-0.0108 (5)	-0.0241 (5)
O2	0.0521 (6)	0.0386 (6)	0.0680 (7)	-0.0148 (5)	-0.0100 (5)	-0.0237 (5)
S1	0.0670 (3)	0.0416 (2)	0.0651 (3)	-0.0161 (2)	-0.0253 (2)	-0.0246 (2)

Geometric parameters (Å, °)

C1—N2	1.3323 (17)	C8—H8	0.9300
C1—N3	1.3367 (19)	C9—N1	1.332 (2)
C1—S1	1.7582 (16)	C9—H9	0.9300
C2—N3	1.329 (2)	C10—C11	1.514 (2)
C2—C3	1.383 (2)	C10—S1	1.7858 (16)
C2—H2	0.9300	C10—H10A	0.9700
C3—C4	1.387 (2)	C10—H10B	0.9700
C3—H3	0.9300	C11—O1	1.1984 (18)
C4—N2	1.3456 (18)	C11—O2	1.3342 (17)
C4—C5	1.4886 (19)	C12—O2	1.4529 (19)
C5—C8	1.385 (2)	C12—C13	1.479 (3)

C5—C6	1.388 (2)	C12—H12A	0.9700
C6—C7	1.381 (2)	C12—H12B	0.9700
C6—H6	0.9300	C13—H13A	0.9600
C7—N1	1.325 (2)	C13—H13B	0.9600
C7—H7	0.9300	C13—H13C	0.9600
C8—C9	1.385 (2)		
N2—C1—N3	127.86 (14)	C11—C10—S1	115.61 (11)
N2—C1—S1	118.93 (11)	C11—C10—H10A	108.4
N3—C1—S1	113.18 (10)	S1—C10—H10A	108.4
N3—C2—C3	123.15 (14)	C11—C10—H10B	108.4
N3—C2—H2	118.4	S1—C10—H10B	108.4
C3—C2—H2	118.4	H10A—C10—H10B	107.4
C2—C3—C4	117.32 (15)	O1—C11—O2	124.33 (14)
C2—C3—H3	121.3	O1—C11—C10	126.61 (14)
C4—C3—H3	121.3	O2—C11—C10	109.02 (13)
N2—C4—C3	120.88 (13)	O2—C12—C13	107.85 (15)
N2—C4—C5	116.22 (12)	O2—C12—H12A	110.1
C3—C4—C5	122.90 (13)	C13—C12—H12A	110.1
C8—C5—C6	116.98 (13)	O2—C12—H12B	110.1
C8—C5—C4	120.72 (13)	C13—C12—H12B	110.1
C6—C5—C4	122.29 (13)	H12A—C12—H12B	108.5
C7—C6—C5	119.32 (15)	C12—C13—H13A	109.5
C7—C6—H6	120.3	C12—C13—H13B	109.5
C5—C6—H6	120.3	H13A—C13—H13B	109.5
N1—C7—C6	124.28 (16)	C12—C13—H13C	109.5
N1—C7—H7	117.9	H13A—C13—H13C	109.5
C6—C7—H7	117.9	H13B—C13—H13C	109.5
C9—C8—C5	119.20 (15)	C7—N1—C9	116.14 (14)
C9—C8—H8	120.4	C1—N2—C4	116.09 (12)
C5—C8—H8	120.4	C2—N3—C1	114.65 (12)
N1—C9—C8	124.07 (17)	C11—O2—C12	116.28 (12)
N1—C9—H9	118.0	C1—S1—C10	101.64 (7)
C8—C9—H9	118.0		

Hydrogen-bond geometry (\AA , $^\circ$)

$D-H\cdots A$	$D-H$	$H\cdots A$	$D\cdots A$	$D-H\cdots A$
C3—H3 \cdots O1 ⁱ	0.93	2.52	3.383 (2)	154
C7—H7 \cdots O1 ⁱⁱ	0.93	2.61	3.373 (2)	140

Symmetry codes: (i) $-x+1, -y, -z+2$; (ii) $x, y, z+1$.